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Inverted encoding models of human population response conflate noise and neural tuning width

Taosheng Liu¹, Dylan Cable² and Justin L. Gardner²

¹Department of Psychology, Michigan State University, East Lansing, MI, USA ²Department of Psychology, Stanford University, Stanford, CA, USA

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Correspondence can be addressed to either Taosheng Liu (tsliu@msu.edu) or Justin Gardner (jlg@stanford.edu).

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3 Taosheng Liu¹, Dylan Cable² and Justin L. Gardner²

4 ¹Department of Psychology, Michigan State University, East Lansing, MI, USA

⁵ ²Department of Psychology, Stanford University, Stanford, CA, USA

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22 Abstract

Channel encoding models offer the ability to bridge different scales of neuronal measurement by 23 interpreting population responses, typically measured with BOLD imaging in humans, as linear sums 24 of groups of neurons (channels) tuned for visual stimulus properties. Inverting these models to form 25 26 predicted channel responses from population measurements in humans seemingly offers the potential to infer neuronal tuning properties. Here, we test the ability to make inferences about neural tuning width 27 from inverted encoding models. We examined contrast invariance of orientation selectivity in human 28 29 V1 (both sexes) and found that inverting the encoding model resulted in channel response functions that became broader with lower contrast, thus, apparently, violating contrast invariance. Simulations 30 showed that this broadening could be explained by contrast-invariant single-unit tuning with the 31 measured decrease in response amplitude at lower contrast. The decrease in response lowers the signal-32 to-noise ratio of population responses that results in poorer population representation of orientation. 33 34 Simulations further showed that increasing signal-to-noise makes channel response functions less 35 sensitive to underlying neural tuning width, and in the limit of zero noise will reconstruct the channel 36 function assumed by the model regardless of the bandwidth of single-units. We conclude that our data are consistent with contrast invariant orientation tuning in human V1. More generally, our results 37 demonstrate that population selectivity measures obtained by encoding models can deviate 38 substantially from the behavior of single-units because they conflate neural tuning width and noise and 39 are therefore better used to estimate the uncertainty of decoded stimulus properties. 40

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42 Significance Statement

43 It is widely recognized that perceptual experience arises from large populations of neurons, rather than a few single-units. Yet, much theory and experiment has examined links between single-units and 44 perception. Encoding models offer a way to bridge this gap by explicitly interpreting population 45 46 activity as the aggregate response of many single neurons with known tuning properties. Here we use this approach to examine contrast invariant orientation tuning of human V1. We show with experiment 47 and modeling that due to lower signal-to-noise, contrast-invariant orientation tuning of single-units 48 49 manifests in population response functions that broaden at lower contrast, rather than remain contrastinvariant. These results highlight the need for explicit quantitative modeling when making a reverse-50 inference from population response profiles to single-unit responses. 51

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54 Introduction

55 Bridging knowledge derived from measurements at different spatial and temporal scales is a significant challenge for understanding the link between neural activity and behavior. While much work has 56 focused on linking single-unit measurements to behavior, there is increasing recognition of the 57 importance of population-scale representations (Benucci et al., 2009; Graf et al., 2011; Churchland et 58 59 al., 2012; Mante et al., 2013; Fusi et al., 2016). In human neuroscience, these bridging challenges are even more severe as many of the core building blocks of knowledge learned from invasive animal 60 experiments are difficult to verify and replicate in humans. It is therefore often unknown whether basic 61 phenomena from the single-unit literature are applicable to humans, let alone how these phenomena 62 will manifest at the larger scale of population activity that is typically interrogated by non-invasive 63 measurement of the human brain. 64

65 Recently, an encoding model approach has proven useful in the analysis of large scale population 66 activity measured by functional imaging (Naselaris et al., 2011; Serences and Saproo, 2012) and offers the promise of bridging knowledge from different species and scales of measurements. Encoding 67 models build off of fundamental results in visual physiology, by encoding complex stimuli in lower 68 dimensional representations such as receptive field or channel models. The assumption is that if these 69 70 neural representations are operative in human cortex, then large-scale measurements of activity represent the aggregated responses of these basic neural operations. For example, a channel encoding 71 model (Brouwer and Heeger, 2009, 2013) has been used to examine continuous stimulus dimensions 72 73 such as color or orientation where it is reasonable to expect that there are large groups of neurons, or channels with known selectivity, and that voxel responses can be modeled as linear combinations of 74 75 such channels. These channel encoding models have been used to examine responses for orientation, color, direction and speed of motion and somatosensory response to better understand apparent motion 76 77 (Chong et al., 2015), cross-orientation suppression (Brouwer and Heeger, 2011), normalization Page 4 / 40

(Brouwer et al., 2015), speeded decision making (Ho et al., 2012), attention (Scolari et al., 2012; Garcia 78 et al., 2013; Saproo and Serences, 2014; Ester et al., 2016), working memory (Ester et al., 2013, 2015), 79 80 perceptual learning (Byers and Serences, 2014; Chen et al., 2015), biases in motion perception (Vintch 81 and Gardner, 2014), and exercise (Bullock et al., 2016) using both functional imaging and EEG (Garcia et al., 2013; Bullock et al., 2016) measurements. Inverting these models to form predictions of channel 82 response from cortical measurements produces tuned response profiles. The interpretations of these 83 84 tuned response profiles are encouraging for the effort of bridging across measurements as they have shown results in concordance with expectations from electrophysiological measurement of phenomena 85 86 such as decision-making reliance on off-target populations (Purushothaman and Bradley, 2005; Scolari et al., 2012) and feature-similarity gain (Treue and Maunsell, 1996; Saproo and Serences, 2014) and 87 response gain (McAdams and Maunsell, 1999; Garcia et al., 2013) modulation effects of attention. 88

89 Here we test the ability of the channel encoding model approach to bridge single-unit and population scale measurement by asking whether the well-known property of contrast-invariant 90 orientation tuning is manifest in predicted channel responses from human primary visual cortex. We 91 92 reasoned that examining whether orientation tuning bandwidth of human cortical population responses 93 change with contrast would provide a good test case for the use of encoding models to bridge 94 measurements, because there is a clear prediction of invariance from single-unit measurements (Sclar and Freeman, 1982). However, contrary to the electrophysiology literature, we found that an encoding 95 96 model produced channel response functions that increased in bandwidth as contrast was lowered. 97 Computational modeling revealed that these effects can be explained by the reduced signal-to-noise ratio of cortical responses at lower contrast. These results emphasize that bridging different levels of 98 99 measurement through these analyses requires explicit quantitative statements of how properties of 100 single-units are expected to manifest in population activity.

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102 Materials and Methods

103 *Subjects*

Six healthy volunteers (ages 33-42, two female) from the RIKEN Brain Science Institute community participated in the experiment; all had normal or corrected-to-normal vision and were experienced subjects in functional imaging experiments. The study protocol was approved by the RIKEN Functional MRI Safety and Ethics Committee and all subjects gave written consent to experimental procedures in advance to participating in the experiment.

109 Stimuli

Stimuli were generated using MGL, a set of Matlab routines for implementing psychophysical experiments (<u>http://gru.stanford.edu/mgl</u>). Stimuli were back projected onto a screen using a LCD projector (Silent Vision 6011; Avotec) at a resolution of 800×600 and a refresh rate of 60 Hz. Subjects viewed the screen via an angled mirror attached to the head coil. The projector was gamma corrected to achieve a linear luminance output.

Visual stimuli were sinusoidal gratings (spatial frequency: 0.7 cpd) in a circular aperture (10°), located to the left or the right of a central fixation cross (1°) at an eccentricity of 8°. The gratings were either low (20%) or high (80%) contrast, and could be in one of eight evenly spaced orientations from 0° to 180° (see Figure 1).

119 Task and Procedures

120 On each trial, two gratings were presented for 5.12 s, followed by a 3.84 s inter-trial interval. 121 During the grating presentation, the phases of both gratings were updated every 0.2 s. The phase of 122 each grating was randomly chosen from one of 16 uniformly distributed phases from 0 to 2π , and the 123 starting time of the phase update of each grating was randomly determined such that the phase updates Page 6 / 40

of the two gratings were asynchronous. The phase updates were implemented to reduce retinal 124 125 adaptation and afterimages. The contrast and orientation of each grating was randomly chosen on each trial such that each combination of contrast (two levels) and orientation (eight levels) was presented 126 three times in each run (48 trials in total). While the inter-trial interval was short which could result in 127 non-linear summation of responses from the previous trial (Boynton, et al., 1996), the trial 128 randomization procedure served to minimize previous trial effects as on average they would come from 129 a random trial type. In addition, a fixation period of 5.12 s preceded each run, making each run 435.2 s 130 in the scanner. Subjects completed 9 runs in the scanner (432 trials in total), which yielded 27 trials per 131 132 orientation/contrast combination.

While the gratings were presented in the periphery, subjects performed a luminance 133 discrimination task at fixation. On each trial in this task, the fixation cross dimmed for 0.4 s twice, 134 separated by a 0.8 s interval, and subjects had to indicate in which interval the cross appeared darker. 135 136 The magnitude of dimming was held constant for one interval while the magnitude of dimming in the 137 other interval was controlled by a one-up two-down staircase. Subjects pressed one of two keys (1 or 2) to indicate their response. The fixation task was performed continuously throughout a run and was 138 asynchronous with the display of the grating stimuli. This task was used to control subjects' attention 139 and ensure a steady behavioral state and eye fixation. The independently randomized contrast and 140 orientations of the two gratings on either side also served as an internal check of the fixation quality, as 141 any systematic bias of eye position for one stimulus would not be systematic for the other. 142

143 MRI methods.

Imaging was performed with a Varian Unity Inova 4T whole-body MRI system (now Agilent Technologies) located at the RIKEN Brain Science Institute, Saitama, Japan. A volume RF coil (transmit) and a four-channel receive array (Nova Medical) were used to acquire both functional and

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Each subject first participated in a separate scanning session to obtain their retinotopic maps (see below for more details), using standard procedures. During this session, a high-resolution 3D anatomical T1-weighted volume (MPRAGE; TR, 13 ms; TI, 500 ms; TE, 7 ms; flip angle, 11°; voxel size, $1 \times 1 \times 1$ mm; matrix, $256 \times 256 \times 180$) was obtained, which served as the reference volume to align all functional images. The reference volume was segmented to generate cortical surfaces using Freesurfer (Dale et al., 1999). Subsequently, the anatomy volumes taken at the beginning of each session were registered to the reference volume so that the cortical regions in the functional scans were aligned with the retinotopy. All analyses were performed in the original (non-transformed) coordinates before being mapped to the cortical surface and specific visual regions.

During the main experiment, functional images were collected using a T2*-weighted echoplanar-imaging sequence (TR, 1.28 s; TE, 25 ms; flip angle, 45°; sensitivity encoding with acceleration factor of 2). We collected 29 slices at an angle approximately perpendicular to the calcarine sulcus, with resolution of $3 \times 3 \times 3$ mm (field of view, 19.2×19.2 cm; matrix size, 64×64). The first four volumes in each run were discarded to allow T1 magnetization to reach steady state. In addition, a T1weighted (MPRAGE; TR, 11 ms; TI, 500 ms; TE, 6 ms; flip angle, 11° ; voxel size, $3 \times 3 \times 3$ mm; matrix, $64 \times 64 \times 64$) anatomical image was acquired to be used for co-registration with the highresolution reference volume collected in the retinotopic session.

Various measures were taken to reduce artifacts in functional images. During scanning, respiration was recorded with a pressure sensor, and heartbeat was recorded with a pulse oximeter. These signals were used to attenuate physiological signals in the imaging time series using retrospective estimation and correction in k space (Hu et al., 1995).

169 *Retinotopic mapping procedure*

In this separate scanning session, we mapped each subject's occipital visual areas using well-170 established phase-encoding methods (Sereno et al., 1995; DeYoe et al., 1996; Engel et al., 1997), so 171 only a brief description is provided here. We presented rotating wedges and expanding/contracting 172 rings over multiple runs and averaged runs of the same type. Then a Fourier analysis was applied to the 173 averaged time course to derive the polar angle map and eccentricity map from the wedge and ring data, 174 respectively. Borders between visual areas were defined as phase reversals in the polar angle map of 175 176 the visual field. The map was visualized on computationally flattened representations of the cortical surface generated by FreeSurfer. For each subject, we could readily define many visual areas, including 177 178 V1, V2, V3, and hV4. However, we will mainly focus on V1 in this study.

179 BOLD Data Analysis.

Data were processed and analyzed using mrTools (http://gru.stanford.edu/mrTools) and other custom code in MATLAB (MathWorks, Natick, MA). Preprocessing of function data included head movement correction, linear detrend, and temporal high-pass filtering at 0.01 Hz. The functional images were then aligned to high-resolution anatomical images for each participant, using an automated robust image registration algorithm (Nestares and Heeger, 2000). Functional data were converted to percentage signal change by dividing the time course of each voxel by its mean signal over a run, and data from the 9 scanning runs were concatenated for subsequent analysis.

187 Voxel selection

We used an event-related (finite impulse response or deconvolution) analysis to select voxels in V1 that responded to the stimulus presentation. Each voxel's time series was fitted with a general linear model with regressors for 16 conditions (2 contrasts × 8 orientations) that modeled the BOLD response in a 25 s window after trial onset. The design matrix was pseudo-inversed and multiplied by the time series to obtain an estimate of the hemodynamic response for each stimulus condition. Because the

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stimulus was independently randomized in the left and right visual field, we fitted two event-related models for each subject, one based on the stimulus in the left visual field and one based on the stimulus in the right visual field.

For each voxel, we also computed a goodness-of-fit measure (r^2 value), which is the amount of 196 variance in the BOLD time series explained by the event-related model (Gardner et al., 2005). In other 197 words, the r^2 value indicates the degree to which a voxel's time course is modulated by the task events, 198 and hence we can use it to select voxels in V1 that were active during the experiment. We selected 199 voxels whose r^2 values were greater than 0.05, which yielded ~100 voxels in each V1 in each 200 hemisphere. Subsequent analysis focused on imaging data in this subset of V1 voxels. Our results did 201 not vary substantially with the voxel selection criterion. For example, when we varied the r^2 cut-off to 202 select a larger number of V1 voxels (~150), the tuning widths of the channel response function were 203 30.6 and 49.9 deg for the high and low contrast stimulus, respectively. For a smaller number of V1 204 voxels (~65), the tuning widths values were 22.7 and 38.7 deg, respectively. 205

206 Channel encoding model

We used a channel encoding model (also referred to as an "encoding model" for brevity below), 207 proposed by Brouwer & Heeger (2009), to characterize the orientation tuning of V1 voxels. 208 Conceptually, the model assumes each voxel's response is some linear combination of a set of 209 210 channels, each channel having the same bandwidth, but with a different preferred stimulus value. We refer to the tuning functions that specify the channels as "model basis functions", which together span 211 the range of all stimulus values. The intuition is that each voxel's response is due to populations of 212 neurons that are tuned to different stimulus values and the analysis proceeds by trying to determine 213 which combination of these neural populations (channels) are most responsible for a voxel's response. 214 215 For every stimulus presentation, the ideal response of each channel is calculated based on the stimulus

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value and model basis functions. The weights of each channel that best fit each voxel's response in the least-squares sense is determined using linear regression from a training data set. Once these weights are fit, the model can be inverted on a left-out test data set to reconstruct channel responses from observed voxel responses. The average channel responses relative to the actual presented stimulus is called a channel response function.

To use as training and test data for the channel encoding model, we obtained single-trial BOLD 221 responses for each V1 voxel with the following procedure. For each V1 hemisphere, we first averaged 222 223 the event-related BOLD response (see voxel selection above) across all voxels and conditions, which 224 served as an estimate of the hemodynamic impulse response function (HIRF) in each V1 hemisphere. We then constructed a boxcar function for each individual trial (with the boxcar length equaling the 225 length of the stimulus presentation), and convolved it with the estimated HIRF, to produce a design 226 matrix coding for each individual trial in each condition. The design matrix was then pseudo-inversed 227 228 and multiplied by the time series to obtain an estimate of the response amplitude for each individual 229 trial in each voxel. We call the set of response amplitudes across all voxels in a V1 hemisphere a response "instance". 230

We fit the encoding model to the instances with a 5-fold cross-validation scheme, in which 4/5231 232 of the trials were randomly selected to be the training data and the remaining 1/5 of trials constituted the test data. This analysis was performed on instances for the high contrast and low contrast trials 233 separately (216 trials per contrast). Our encoding model consisted of 8 evenly-spaced channels (i.e., 234 model basis functions) from 0 to 180°, with each channel a half-wave rectified sinusoid raised to the 235 power of 7. These basis functions were chosen to approximate single-neuron's orientation tuning 236 237 function in V1. In the following exposition, we adopted the notation from Brouwer & Heeger (2009). The training instances can be expressed as a $m \times n$ matrix **B**₁, where m is the number of voxels, and n is 238 number of trials in the training data. We then constructed hypothetical channel outputs given the 239 Page 11 / 40

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stimulus orientation on each trial of the training dataset, which yielded a $k \times n$ matrix C₁, where k is the number of channels (i.e., k=8). Each column in the C₁ matrix represented a set of ideal response to the stimulus orientation on that trial from the eight channels. A weight matrix W ($m \times k$) relates the observed data B and the hypothetical responses

$$\mathbf{B}_1 = \mathbf{W} \mathbf{C}_1 \tag{Equation 1}$$

Each row of **W** represents the relative contribution of the eight channels to that voxel's response. The least-square estimate of **W** was obtained with the following equation (T indicates the transpose of the matrix):

$$\hat{\mathbf{W}} = \mathbf{B}_{1} \mathbf{C}_{1}^{\mathrm{T}} (\mathbf{C}_{1} \mathbf{C}_{1}^{\mathrm{T}})^{-1}$$
(Equation 2)

The test instances can be expressed as a $m \times p$ matrix **B**₂, where *p* is the number of trials in the test data. The estimated channel response to each test stimulus (\hat{C}_2) can then be estimated using the weights **W**:

$$\hat{\mathbf{C}}_{2} = (\hat{\mathbf{W}}^{\mathrm{T}} \hat{\mathbf{W}})^{-1} \hat{\mathbf{W}}^{\mathrm{T}} \mathbf{B}_{2}$$
(Equation 3)

 \hat{C}_2 is a $k \times p$ matrix, with each column representing each channel's response to the stimulus on 253 that test trial. The columns of \hat{C}_2 were circularly shifted such that the channel aligned to the test 254 stimulus on that trial was centered in the orientation space. The shifted columns were then averaged to 255 obtain a mean channel response. This procedure was repeated in each fold of the cross-validation, and 256 the mean channel responses from each fold were further averaged to obtain what we will refer to as a 257 258 "channel response function". For each V1 hemisphere in each participant, we obtained channel response functions for both the contralateral and ipsilateral stimulus, separately for the high contrast 259 and low contrast conditions. 260

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In addition to the channel response function, we also calculated a goodness-of-fit measure of Page 12 / 40 267

the encoding model. In each cross-validation fold, after we obtained estimates of the channel weights \hat{W} , we also constructed another set of hypothetical channel outputs given the actual stimulus orientation in the test data, C₂ (note that this is a matrix similar to C₁ but for the test trials, and differs from \hat{C}_2 which is the *estimated* channel responses based on the voxel responses). The predicted voxel response can be obtained via:

$$\mathbf{B}_{2\mathsf{pred}} = \mathbf{\hat{W}} \mathbf{C}_2 \tag{Equation 4}$$

To the extent that the encoding model provides a good fit to the data, the predicted response B_{2pred} ($m \times p$) should be similar to the observed response, B_2 . We can thus calculate the amount of variance explained by the model as

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$$r^2 = 1 - \frac{\sum (B_{2pred} - B_2)^2}{\sum (B_2 - \bar{B}_2)^2}$$
 (Equation 5)

\overline{B_2} is the mean voxel responses across voxels and trials, and the summation was performed across voxels and trials (i.e., all values in the $m \times p$ matrix). We calculated r² from each fold of the cross validation and averaged them across the folds to obtain a single measure of the goodness-of-fit, for each contrast. Note this r² is different from the r² that represents the goodness-of-fit of the event-related model (see above section *Voxel selection*). In the remainder of this report, we will focus on this r² value that indexes the goodness-of-fit of the channel encoding model.

278 *Quantifying the channel response function*

279 We fitted a circular bell shaped function (von Mises) to the channel response function

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$$y = y_0 + Ae^{\kappa cos(x-\mu)}$$
(Equation 6)

Where κ is the concentration parameter which controls the width of the function, μ is the mean, *x* is the orientation, *A* is the amplitude parameter, and y_0 is the baseline. Thus there were four free parameters of the fit: baseline (y_0), amplitude (*A*), mean (μ), and concentration (κ). Because orientation is on [0 π], Page 13 / 40 whereas the von Mises function spans the interval $[0 \ 2\pi]$, orientation values were multiplied by 2 during the fit, after which the fitting results were scaled back to $[0 \ \pi]$. Fitting was performed using a non-linear least square method, as implemented in MATLAB. To ease the interpretation of the results, we report the half-width-at-half-maximum instead of the concentration parameter, κ , because the latter is inversely related to the variance.

289 Linking neuronal tuning to channel response: a computational model

290 We implemented a computational model that links neuronal tuning to channel response functions, using the assumptions underlying the channel encoding model. Given the channel response 291 function is a highly derived statistic, this modeling effort was used to clarify how various assumptions 292 293 of neural tuning and signal-to-noise would manifest in channel response functions. Specifically, we used this model to fit our observed data and simulate other scenarios to test the validity of the encoding 294 295 model. The schematic of the model is outlined in Figure 3. The model contains 100 V1 voxels, 296 comparable to the actual number of voxels used in our data analysis (see above). Each voxel is assumed 297 to contain neurons tuned to all orientations, whose tuning functions are described by von Mises functions (see above). The preferred orientation (μ) are evenly distributed across all possible 298 orientations in 1° increment (left column, "Neural tuning functions"), forming 180 classes of neurons. 299 300 The width of the neural tuning function is specified by the concentration parameter, κ , which is the same for all neurons but can be manipulated across different simulations. The area under the orientation 301 tuning function for neurons was normalized to one so that average firing rate for each neuron would not 302 303 vary with tuning width. For each voxel, a weight vector, w, was generated by randomly sampling 180 numbers from [0 1]. This weight vector specifies how much each class of neurons contributes to the 304 305 voxel's response. For an input stimulus with an orientation, θ , the response of each neuron is calculated according to its tuning function (see Eq. 6). For each voxel, the responses of individual neurons are 306 multiplied by the weight vector, w, and then summed to arrive at a predicted response (middle column, 307 Page 14 / 40

³⁰⁸ "Random weights"). To calculate the final voxel response, Gaussian noise with $N(0, \sigma)$, is then added ³⁰⁹ to this response to simulate physiological and thermal noise in BOLD measurements. Thus, each ³¹⁰ voxel's response is determined by neuronal tuning width (κ), weight vector (w), and noise (σ). Note w³¹¹ is randomly generated for each voxel in each simulation, whereas κ and σ are parameters that we ³¹² examined systematically in several simulations (see Results).

We simulated experiments with the same basic set-up as our empirical study: 8 possible orientations, with each orientation shown 27 times (trials). For each trial, we obtained a vector of voxel responses (instances), calculated as above. Then all the trial instances were subjected to the same analysis as the real data described above, i.e., cross-validation in which 4/5 of data were used to obtain the channel weights and 1/5 of the data were used to obtain channel response functions. We used the same exact code to analyze the synthetic and real data.

319 *Computation of the posterior distributions*

We computed the probability of different stimulus values given the test data, i.e., the posterior distribution, using a technique from van Bergen et al. (2015). The method begins with finding the weights of the channel encoding model as above. After removing the signal due to the encoding model, a noise model is fit to the residual response. The noise model assumes that each individual voxel's variability is Gaussian with one component which is independent among voxels and another that is shared across all voxels. Each of the channels is modeled to have independent, identically distributed Gaussian noise. This leads to a covariance matrix for the noise as follows:

$$\mathbf{\Omega} = \rho \mathbf{\tau} \mathbf{\tau}^{\mathsf{T}} + (1 - \rho) \mathbf{I} \bullet \mathbf{\tau} \mathbf{\tau}^{\mathsf{T}} + \sigma^2 \mathbf{\hat{W}} \mathbf{\hat{W}}^{\mathsf{T}}$$
(Equation 7)

Where Ω is the noise model's covariance matrix, I is the identity matrix, \bullet denotes element-wise multiplication, τ is a vector containing each voxel's independent standard deviation, ρ is a scalar between 0 and 1 which controls the amount of shared variability among voxels, σ is the standard

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deviation of each channel and $\hat{\mathbf{W}}$ is the estimated weight matrix from Equation 2. This noise model 331 with parameters, τ , ρ and σ is fit via maximum likelihood estimation to the residual and can be used to 332 compute the probability of generating any particular response given a stimulus value. Inversion of this 333 334 equation using Bayes' rule and a flat prior allows one to compute the probability of any stimulus value given a response-the posterior distribution. For a derivation and detailed explanation see (van Bergen 335 et al., 2015). All analysis followed the same 5-fold cross-validation scheme used for the encoding 336 model by which 4/5 of the data were used to fit the model weights and noise model parameters and the 337 338 1/5 left out data were used to compute the posterior distribution using Bayes' rule. Results are shown 339 averaged across all five left-out folds.

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341 Results

We measured BOLD responses from retinotopically defined V1 to oriented sinusoidal gratings 342 (Fig 1) and used the resulting data to train and test a channel encoding model. We first report average 343 (univariate) activity across voxels. While responses averaged across subjects and voxels in V1 did not 344 345 show any apparent selectivity for orientation (Fig 2A), they were, as expected (Boynton et al., 2012), greater for higher contrast compared to low contrast contralateral stimuli (compare red vs. yellow). 346 Consistent with the laterality of V1, responses did not vary with contrast of the ipsilateral stimulus 347 (black). The presence of a strong contrast response for the contralateral stimuli but a complete absence 348 349 of such a response for ipsilateral stimuli also suggest that subjects maintained stable central fixation.

Despite this lack of orientation selectivity, channel response functions obtained from the 350 encoding model and averaged across subjects displayed peaked responses at the true orientation of the 351 352 contralateral stimulus for both high (Fig 2B, red) and low (Fig 2C, yellow) contrast stimuli. We computed r², a measure of goodness-of-fit, which showed that the channel encoding model accounted 353 for 31% and 13% (high and low contrast contralateral stimuli, respectively) amount of the variability of 354 the data. The ability to recover these peaked function of orientation is consistent with previous studies 355 using classification approaches (Kamitani and Tong, 2005) and is presumably due to biases in response 356 357 to orientation which differ for individual voxels but are eliminated when responses are averaged across voxels. As expected from the lateralization of V1, these channel response functions are flat when 358 constructed for the ipsilateral stimulus (black), thus serving as an internal control on the validity of the 359 encoding model approach. 360

However, contrary to the expectations of contrast-invariance, channel response functions were broader for low contrast compared to high contrast contralateral stimuli (Fig 2D, compare yellow to red). Fitting a bell shaped function (von Mises) to the subject-averaged channel response functions

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revealed that the tuning width went from 25.6 to 42.0 deg (half-width at half-height) as contrast was 364 365 lowered. Amplitude was also decreased for the low contrast stimuli from 0.27 to 0.23, where 1 would be the ideal height of the channel response function if voxel responses contained noise-free information 366 about stimulus orientation. This pattern of results was also evident in individual subject's channel 367 response functions. 9 out of 12 hemispheres showed a decrease in tuning width as contrast was 368 decreased (p = 0.013, t(11) = -2.60, one-tailed paired t-test). 10 out of 12 hemispheres had lower 369 amplitude for the low contrast condition (p = 0.042, t(11) = 1.90, one-tailed paired t-test). We also 370 examined extrastriate areas V2, V3, and hV4 and found similar results. In V2, tuning width went from 371 372 21.3 to 75.5 deg and amplitude went from 0.25 to 0.19 as contrast was lowered. In V3, tuning width went from 17.8 to 88.6 deg and amplitude went from 0.23 to 0.18 as contrast was lowered. Finally, in 373 374 hV4, tuning width for high contrast stimuli was 14.2 deg and channel response function was essentially 375 flat for low contrast stimuli. Below we will focus on results from V1, which is best informed by neurophysiological results (Sclar and Freeman, 1982; Skottun et al., 1987; Carandini et al., 1997). 376

377 While the lack of contrast-invariant channel response functions might imply broader neuronal tuning at low contrast in human visual cortex, we instead considered whether it might be due to the 378 weaker stimulus-driven signal at lower contrast. As noted above, BOLD responses had lower amplitude 379 with lower contrast (Fig 2A). Given that many sources of noise in BOLD are non-neural (e.g. 380 hemodynamic variability and head motion) and thus not expected to vary with signal strength, these 381 lower amplitude responses result in lower signal-to-noise ratio (SNR) of the measurements made with 382 383 lower contrast. Indeed, the amount of variance accounted for by the encoding model, r^2 , was 384 significantly lower for low contrast compared to high contrast stimuli for 12 out 12 hemispheres (p \leq 0.001, t(11) = 5.58, one-tailed paired t-test). In the extreme case, channel response functions built on 385 responses without any signal, as for the ipsilateral stimulus, are flat. Thus, we reasoned that the lower 386 SNR measurements at low contrast could also result in flatter, i.e. broader, channel response functions. 387

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To test whether reduced SNR at low contrast, rather than changes in neural tuning, could 388 account for the increased channel response bandwidth at low contrast, we built simulations (Fig 3, see 389 Materials and Methods). Briefly, each simulated voxel received randomly weighted responses from 390 simulated orientation-tuned neurons. Different voxels had different weightings of the neuronal 391 responses, thus resulting in weak, but different orientation selectivity across voxels. We added random 392 Gaussian noise to these voxel responses and trained and tested the encoding model using the same 393 394 procedures as we did for the actual BOLD data. We varied the standard deviation of the added noise (σ) to produce channel response functions that best fit the empirical data (in the least-squares sense) 395 396 from the high contrast trials (Fig 4A). We then noted that in the empirical data there was a 42.2% decrease in neural response from high to low contrast across voxels (Fig 2A). We therefore decreased 397 398 neural response by this amount for all the simulated neurons and found that the resulting channel 399 response function to be a reasonable fit for the low contrast data (Fig 4B). Importantly, this good correspondence between model predictions and data was achieved without fitting any parameter, 400 because the only thing we changed in the simulation was to decrease the magnitude of response across 401 402 all neurons according to the value found from the empirical data. This suggests that reductions in response magnitude, and therefore SNR, are sufficient to produce changes in channel response width 403 404 commensurate with what we observed.

While tuning width is not expected to change with contrast, what if we had tested a property for which we expect a neural tuning width change? Do channel response functions track changes in neural tuning width? We simulated neural tuning functions from 5 to 40 degrees half-width at half-height as well as a "stick function" which responds only to a single orientation maximally and does not respond to any other orientation and computed channel response functions under different amounts of noise (Fig 5A, cyan to magenta curves represent different neural tuning widths). We found that the resulting channel response functions did indeed track the neural tuning widths, but as the goodness-of-fit r^2 ,

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412 increased (achieved by varying the standard deviation of the added noise, abscissa in Fig 5B), the 413 difference in channel response widths was diminished (note the larger splay of curves on the left vs 414 right side of Fig 5B). Thus, channel response functions can reflect underlying neural tuning widths, but, 415 perhaps counterintuitively, the better the goodness-of-fit of the encoding model, the less difference the 416 neural tuning width makes.

To understand why better goodness-of-fit implies worse model discriminability of underlying 417 neural tuning functions, it is important to note that the absolute width of channel response functions is, 418 419 in the limit of no noise, determined by the basis functions used in the encoding model and not by the 420 neural tuning widths themselves. We simulated two variations of the encoding model in which we varied the model basis function widths. We increased the basis function width by decreasing the 421 exponent on the sinusoidal basis to 3 (Fig 5A) and decreased the basis function width by using stick 422 filters (Fig 5C). For all the simulations, as the goodness-of-fit increases, the recovered channel 423 424 response functions approach the width of the model basis function (dashed line) rather than the 425 neuronal response width (note that for both the channel response functions and the model basis functions, we use the fitted half-width at half-height as our measure of tuning width and therefore the 426 stick functions do not have infinitely narrow tuning). Given that the encoding model is essentially a 427 428 linear regression model, this is not an unexpected outcome. Linear weights are being determined to best map the voxel responses into ideal channel responses. As long as the voxel responses are determined 429 by stimulus orientation, then the regression model will be able to recreate exactly any model basis 430 431 functions that can be formed as linear combinations of the represented orientations

The above analysis suggests that while absolute neural tuning width may not be readily determined from the channel response function width, *changes* in tuning width might be meaningful if SNR does not change between conditions. That is, reading vertically for one level of goodness-of-fit in Fig 5A-C, the channel response function width changes systematically as a function of neural tuning Page 20 / 40 width. Might we be able to determine that neural tuning width has changed if we observe data in which
we have matched goodness-of-fit of the encoding model? While this does not occur for changes in
contrast, this could be the case for potential modulation of tuning width by cognitive factors like
attention and learning.

440 However, changes in neural tuning width also result in changes in the model's goodness-of-fit thus complicating the possibility of interpreting changes in channel response width. We simulated 441 different neural tuning widths from 5-45 degrees half-width at half-height (Fig 6, abscissa) for different 442 443 amounts of additive noise (standard deviation ranging on a logarithmic scale from 0.001 to 1, blue to vellow curves). As can be appreciated by the downward slopes of the curves in Fig 6, as the neural 444 tuning widths get larger, the fit of the encoding model gets worse (lower r^2 , ordinate). The reason for 445 this worsened fit is because as neural tuning width gets wider, there is less information about 446 orientation available to fit the model. In the extreme, a flat neural tuning function would result in no 447 448 orientation specific response and the encoding model would fail to fit the data completely. Each curve 449 in the simulation is what one might expect to measure if neural tuning width is the only variable that changes in the experiment and noise is due mostly to external factors that do not change with 450 conditions. That is, if one expects only a change in neural tuning width, the resulting channel response 451 functions would be expected to have both wider tuning and lower r^2 . This pattern of results would make 452 it difficult, if not impossible, to determine whether the changes in tuning were due to decreased SNR, 453 decreased neural tuning width, or some combination of both. 454

These empirical and simulation results suggest caution in interpreting changes in channel response functions because they could be due to either or both changes in neural tuning width and changes in signal strength. What better ways are there for interpreting these functions? A recent report (van Bergen et al., 2015) proposes to transform these channel response functions into posterior distributions that show the probability of different stimulus values given the measured response. We Page 21 / 40

applied the same analysis to our channel response functions, by estimating the distribution of noise in 460 the voxels and model to determine the probability of measuring responses given any oriented stimulus 461 and then applying Bayes' rule with a flat prior to obtain the probability of various stimuli given the 462 463 responses we measured in a left-out validation set of data (see Materials and Methods for details, Fig 7). This resulted in posterior distributions that were peaked around the actual orientation for the 464 contralateral stimulus (red and orange curves), but flat for the ipsilateral stimulus (black curves), thus 465 replicating the results on channel responses. This transformation of the results into posterior 466 distributions allows for a more straight-forward interpretation of the encoding model approach — 467 468 encouraging interpretation in terms of the certainty by which a given neural response tells us what the stimulus was, rather than what it implies about underlying neural tuning functions. 469

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472 Discussion

Using an encoding model approach we built channel response functions for orientation and found that, 473 474 unlike contrast-invariant single-units, they were broader at lower contrast. Simulations showed that this 475 effect could be fully explained by the measured decrease in overall neural response between high and low contrast, which results in lower signal-to-noise ratio. As signal-to-noise is increased, channel 476 response functions become narrower, until, in the limit of no noise, they approximate the shape of the 477 478 model basis function (and not the underlying neural tuning function). While changes in underlying 479 neural selectivity in our model could be reflected in channel response functions, our results demonstrate that changes in channel response functions do not necessarily reflect changes in 480 underlying neural selectivity. 481

Orientation selectivity of single-unit responses have been shown to be invariant to image 482 483 contrast (Sclar and Freeman, 1982; Skottun et al., 1987; Carandini et al., 1997), suggestive of a general neural computational mechanism (Carandini and Heeger, 2012) by which visual perception can remain 484 relatively unaffected by differences in visibility of stimuli. Despite this central theoretic importance, 485 486 obtaining non-invasive measurement of selectivity bandwidth from human cortex has been technically difficult because orientation selectivity is organized into cortical columns (Hubel and Wiesel, 1962, 487 1968; Blasdel and Salama, 1986; Bonhoeffer and Grinvald, 1991), much smaller than the typical spatial 488 489 resolution of blood-oxygen level dependent (Ogawa et al., 1990, 1992) measurements. While direct measurements of such columnar structures in humans has been achieved (Cheng et al., 2001; Sun et al., 490 491 2007; Yacoub et al., 2007, 2008), multivariate analysis using pattern classification approach to decode orientation and motion direction (Haynes and Rees, 2005; Kamitani and Tong, 2005, 2006) from 492 distributed activity patterns has become a more common approach (Norman et al., 2006). However, this 493 494 classification approach generally produces a categorical outcome, for example, which of two 495 orientations was more likely to have resulted in the measured response pattern and thus is not typically Page 23 / 40

used for probing selectivity bandwidth of neural representations. The encoding model approach allows
one to reconstruct a response profile for a stimulus that has a tuning bandwidth that can be inspected
across different contrasts.

499 While we found an increase in the bandwidth of channel response functions for lower contrast 500 stimuli from human V1, this increase could be fully accounted for by the measured reduction of response amplitude due to contrast, thus reconciling our data with contrast-invariant orientation tuning. 501 We recognize that contrast invariance at the population level as measured with BOLD is not guaranteed 502 503 even if single-unit spiking responses display contrast-invariance. Systematic relationships between 504 contrast-sensitivity and selectivity for orientation could result in population responses changing selectivity with contrast. For example, if the least orientation-selective neurons saturate their responses 505 at lower contrast than the most selective neurons, then population response would become more 506 selective as contrast increases because population response would be dominated by the most selective 507 508 neurons. However, no such systematic relationship has been observed and population spiking responses 509 appear contrast-invariant in cats (Busse et al., 2009), consistent with our results. Furthermore, BOLD 510 measurements may be better correlated with local field potentials than spiking activity (Logothetis et al., 2001), which could also result in deviations of BOLD population measures of contrast invariance 511 and spiking activity of neurons. If BOLD measures are sensitive to sub-threshold, synaptic activity that 512 can contribute to local field potentials, broadening of channel response functions that we observed 513 could be reflective of sub-threshold activity, if such activity is not contrast-invariant. However, 514 515 intracellular measurements of membrane potentials show that selectivity does not broaden at lower 516 contrast, in fact, selectivity is slightly increased at low contrast (Finn et al., 2007), consistent with our interpretation that channel response function broadening at low contrast is due to reduction in signal-to-517 noise. 518

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Given our results, bridging effects of attention on single units with effects uncovered using Page 24 / 40

encoding models of functional imaging measurements (Sprague et al., 2015) may be similarly 520 complicated as bridging contrast invariance effects. Single unit studies have suggested that neurons 521 change gain, not selectivity bandwidth (McAdams and Maunsell, 1999; David et al., 2008) with spatial 522 523 attention, a key finding that has shaped our understanding of neural mechanisms of attention (Carrasco, 2011; Ling et al., 2015). In human population measurements, improved orientation encoding has been 524 found when orientation (but not contrast) is task relevant (Jehee et al., 2011; Ling et al., 2015). While it 525 would be of interest to know whether these population effects of attention reflect differences in neural 526 tuning bandwidth, selective attention, like image contrast, also modulates response amplitudes 527 528 (Brefczynski and DeYoe, 1999; Gandhi et al., 1999; Kastner et al., 1999; Kastner and Ungerleider, 2000; Reynolds and Chelazzi, 2004) and thus is expected to improve signal-to-noise ratio for 529 530 population measures. Similarly to contrast effects, attention should be expected to bias channel 531 response functions toward a narrower tuning even if neural tuning bandwidth does not change.

A similar disconnect between single-unit and population measures impacts even simpler 532 measures of cortical response that do not require multivariate approaches. Contrast sensitivity can be 533 directly imaged because single-units monotonically increase response with contrast (Albrecht and 534 Hamilton, 1982; Sclar et al., 1990; Busse et al., 2009) resulting in a population response that also 535 monotonically increases (Tootell et al., 1998; Boynton et al., 1999; Logothetis et al., 2001; Avidan and 536 Behrmann, 2002; Olman et al., 2004; Gardner et al., 2005). Spatial attention has generally been shown 537 to shift contrast response vertically upward when measured with functional imaging (Buracas and 538 539 Boynton, 2007; Li et al., 2008; Murray, 2008; Pestilli et al., 2011; Hara and Gardner, 2014), which 540 appears to be different from the variety of effects from contrast-gain to response-gain reported for single-units (Reynolds et al., 2000; Martínez-Trujillo and Treue, 2002; Williford and Maunsell, 2006; 541 Lee and Maunsell, 2010; Pooresmaeili et al., 2010; Sani et al., 2017). Consideration of normalization 542 and the size of the attention field relative to stimulus-driven responses can give rise to effects that can 543

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account for single-unit responses and EEG measures (Reynolds and Heeger, 2009; Itthipuripat et al., 2014). But, predictions of this normalization model of attention may differ for single-units and population measures as different neurons in a population can be exposed to different balance of attention field and stimulus-drive, giving rise to additive shifts when considered as a population (Hara et al., 2014). Relatedly, response gain changes may also manifest as additive shifts when directly examining voxel feature selectivity (Saproo and Serences, 2010).

While neural tuning width can be reflected in channel response functions, neural tuning width 550 551 and signal-to-noise changes are intertwined, making it hard to disentangle their effects. For example, 552 one might examine conditions in which signal-to-noise is matched and then hope to attribute changes in channel response function bandwidth solely to changes in neural tuning bandwidth. However, our 553 simulations show signal-to-noise measures such as the variance accounted for by the encoding model 554 (r^2) , covaries with neural tuning width. As neural tuning width broadens there is less modulation of 555 voxel response with orientation and thus the encoding model shows a decrease in r^2 . Therefore, even 556 pure changes in neural tuning width would result in conditions with lower r², making it hard to attribute 557 changes in channel response functions solely to changes in neural tuning width. 558

559 The results of our simulation are agnostic to the source of selectivity for orientation in voxels. 560 One possible source of orientation information are the irregularities of columnar organization which could give rise to small, idiosyncratic biases in voxels (Boynton, 2005; Swisher et al., 2010). However, 561 large scale biases for cardinal (Furmanski and Engel, 2000; Sun et al., 2013) and radial (Sasaki et al., 562 2006) orientations have been reported, and these biases have been shown to be an important source of 563 information to drive classification (Freeman et al., 2011, 2013; Beckett et al., 2012; Wang et al., 2014; 564 565 Larsson et al., 2016), but see (Alink et al., 2013; Pratte et al., 2016). Large-scale biases may result from vascular (Gardner, 2010; Kriegeskorte et al., 2010; Shmuel et al., 2010) or stimulus aperture (Carlson, 566 2014) related effects. Our simulations do not require, or exclude, any topographic arrangement of 567 Page 26 / 40

biases. Regardless of the source of orientation bias, channel response function widths would beexpected to broaden as signal-to-noise decreases.

More generally, our results suggest a "reverse-inference" problem (Aguirre, 2003; Poldrack, 570 2006) when interpreting outputs from inverted encoding models. Forward encoding from hypothetical 571 572 neural responses to population activity is a powerful tool, but reversing this process to infer about neural responses is problematic when there is not a one-to-one mapping between single-unit and 573 population measures. Consequently, this reverse-inference problem is not restricted to channel 574 575 encoding models, but will occur for other encoding model approaches such as population receptive 576 fields (Dumoulin and Wandell, 2008) or Gabor wavelet pyramids (Kay et al., 2008), if one were to invert these models to infer properties of the underlying neural responses. For contrast and orientation, 577 both increases in response amplitude and neural selectivity can result in narrower bandwidth of the 578 channel response functions so reverse inference requires taking both into account. Regardless of which 579 580 neural change has occurred, read-out of these responses, be they in the brain or from external 581 measurement, will have less certainty about what stimulus has caused those responses. Techniques that 582 represent the output of encoding models as posterior distributions (van Bergen et al., 2015) offer a straightforward interpretation of the uncertainty in determining stimulus properties from cortical 583 584 responses.

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Figure 1. Schematics of the experiment. Low and high contrast gratings of eight possible orientations were presented, with contrast and orientation independently randomized for each visual field. A luminance change-detection task was performed at the fixation to control subjects' fixation and state of attention.

Figure 2. (A) Group-averaged mean BOLD response across V1 for each orientation, separately for the 591 contralateral and ipsilateral high- and low- contrast stimuli. (B) Group-averaged channel response 592 functions from V1 to a high contrast grating, contra: response calculated for the contralateral stimulus; 593 ipsi: response calculated for the ipsilateral stimulus. (C) Same as B except for low contrast grating. (D) 594 Group-averaged channel response functions to the low and high contrast grating in the contralateral 595 596 visual field (symbols, same as the contralateral response in B and C). Solid lines are best fitting von Mises functions to each contrast level. Error bars in these graphs are standard error of the mean (s.e.m.) 597 across participants. 598

Figure 3. Schematic of the model linking neuronal response to channel response. Each voxel (right column) received randomly weighted responses from orientation-tuned neurons (left column). After weighting and summing, random Gaussian noise was added to obtain simulated voxel responses (see text for more details).

Figure 4. Model predictions of empirical channel response functions. (A) Empirical channel response function for contralateral high contrast stimuli (red symbols, same data as in Fig 2B) were fit by the computational model, with the best-fitting channel response shown in black symbols and lines. (B) Empirical channel response function for contralateral low contrast stimuli (red symbols, same data as in Fig 2C) and the channel response from the same model used in A (black symbols and lines), except that the neuronal response amplitude was reduced. Error bars are standard error across subjects and Page 28 / 40

609 hemispheres.

Figure 5. Model simulations of how channel response function varies with neural tuning, signal-tonoise ratio, and model basis function. Each panel uses a different model basis function (shown in the top graph) to derive channel responses from synthetic data generated with different combinations of signal-to-noise (r^2 , x-axis) and neural tuning width (colored lines). The width of the channel response function is plotted on the y-axis. Horizontal dashed lines indicate the width of the model basis function.

Figure 6. Model simulations of how goodness-of-fit of the encoding model (r^2) varies with neural tuning width and noise level in the synthetic data. Different colors represent different amount of Gaussian noise added to the simulated neural response.

Figure 7. Posterior distributions from the Bayesian analysis. These functions represent the probability that a given stimulus (measured by the offset from the true orientation, x-axis) caused the observed BOLD response. Posterior distributions for high and low contrast contralateral stimuli are shown in red and yellow, respectively, whereas posterior distributions for ipsilateral stimuli are shown in black. Shaded region represents the standard error over subjects and hemispheres.

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